

ANNOTATED SHEET

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+1 TCACTAGTACCATAATATTACAAAACACACACCTCAGAAACGAAGCTTGC  
+53 ACATAATGTCTCTTCAACAAGTACCCACAACAGAAACAACCACCAGGAAGAA  
1 M S L Q Q V P T T E T T T R K N  
+105 CCGAAACGAGGGGCGGAAACGATTTACCGACAAACAAATAAGTTTCCTAGAG  
17 R N E G R K R F T D K Q I S F L E  
+157 TACATGTTTGTAGACACAGTCGAGACCCGAGTTAAGGATGAAACACCAGTTGG  
34 Y M F E T Q S R P E L R M K H Q L  
+209 CACATAAACTCGGGCTTCATCCTCGTCAAGTGGCGATATGGTTCCAGAACAA  
51 A H K L G L H P R Q V A I W F Q N K  
+261 ACGCGCGCGATCAAAGTCGAGGCAGATTGAGCAAGAGTATAACGCGCTAAAG  
69 R A R S K S R Q I E Q E Y N A L K  
+313 CATAACTACGAGACGCTTGCGTCTAAATCCGAGTCTCTAAAGAAAGAGAATC  
86 H N Y E T L A S K S E S L K K E N  
+365 AGGCCCTACTCAATCAGgtatggttgcaaacttacaatgttgcatccaacta  
103 Q A L L N Q  
+417 ttttaagtagttttgaattttgtgacaataaagattgacaaatgttggtttga  
+469 taattgattaacagTTGGAGGTGCTGAGAAATGTAGCAGAAAAGCATCAAGA  
109 L E V L R N V A E K H Q E  
+521 GAAAACTAGTAGTAGTGGCAGCGGTGAAGAATCGGATGATCGGTTTACGAAC  
122 K T S S S G S G E E S D D R F T N  
+573 TCTCCGGACGTTATGTTTGGTCAAGAAATGAATGTTCCGTTTTCGCGACGGTT  
139 S P D V M F G Q E M N V P F C D G  
+625 TTGCGTACTTTGAAGAAGGAAACAGTTTGTGAGATTGAAGAACAACCTGCC  
156 F A Y F E E G N S L L E I E E Q L P  
+677 AGACCCTCAAAAGTGGTGGGAGTTCTAAAGAGTAAAGAAGGATGTAGAAGTA  
174 D P Q K W W E F \*  
+729 GTAGAGTAAAACTAAAACATACCAGATAGTTGGTTTACACTTTGT

Athb-1 LPEKRRRLTTEQVHLLKSFETENKLEPERKTQLAKKLGLOPROVAVWFQNRARWTKQLERDYDLKSTYDQLLSNYDSIVMDNKLSEVTSLTEK  
Athb-6 LSEKRRRLSINQVKALEKNFELENKLEPERKVLAQELGLOPROVAVWFQNRARWTKQLEKDYGVLTQYDSLRRHFDLSRRDNESLLQEISKLTKE  
Athb-7 NKNNQRRFSDQIKSLEMMFSETRLEPRKKVOLARELGLOPROVAIWQNKRRWKSQLETEYNILRQYDNLASQFESLKKEQALVSELQRLKEA  
Athb-12 KSNQKRFNEEQIKSLELIFESETRLEPRKKVOVARELGLOPRQMTIWFQNKRRWTKQLEKEYNTLRANYNNLASQFEIMKKEQSLVSELQRLNEE  
Hahb-4 RNEGRKRFTRQISFLEYMFETQSRPELRMKHQLAHKLGLOPROVAIWQNKRRSKSRQIEQYNAKKNYETLASKSESLLKKNQALLNQLVLRNV

FIG. 1

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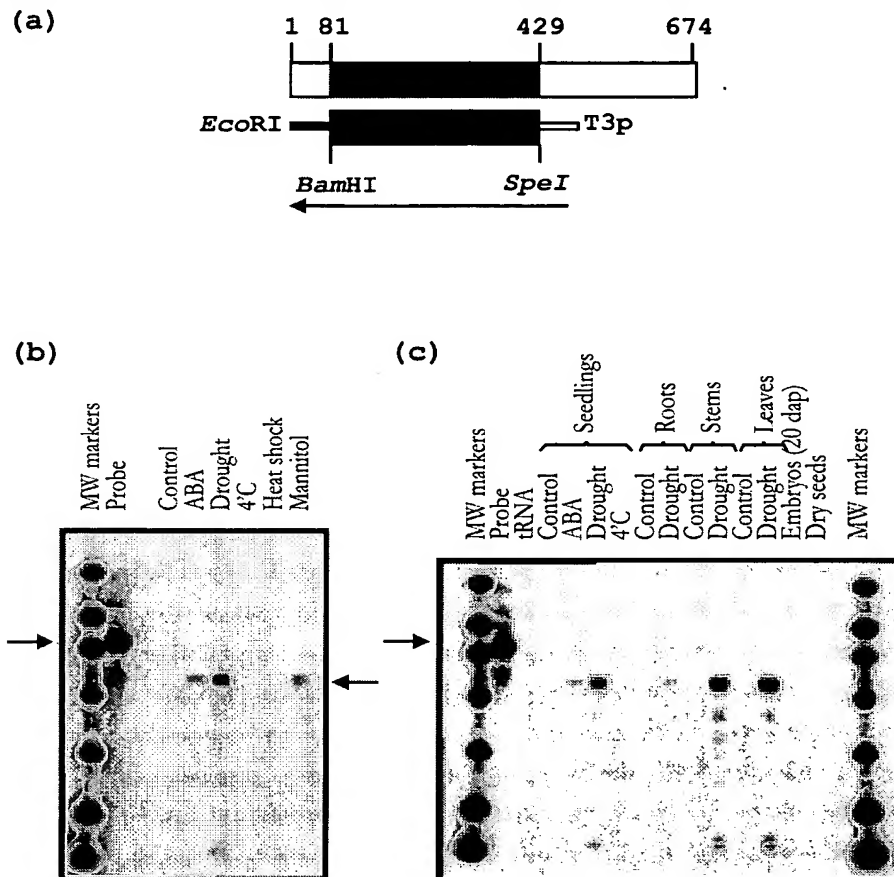


FIG. 2

ANNOTATED SHEET



FIG. 22

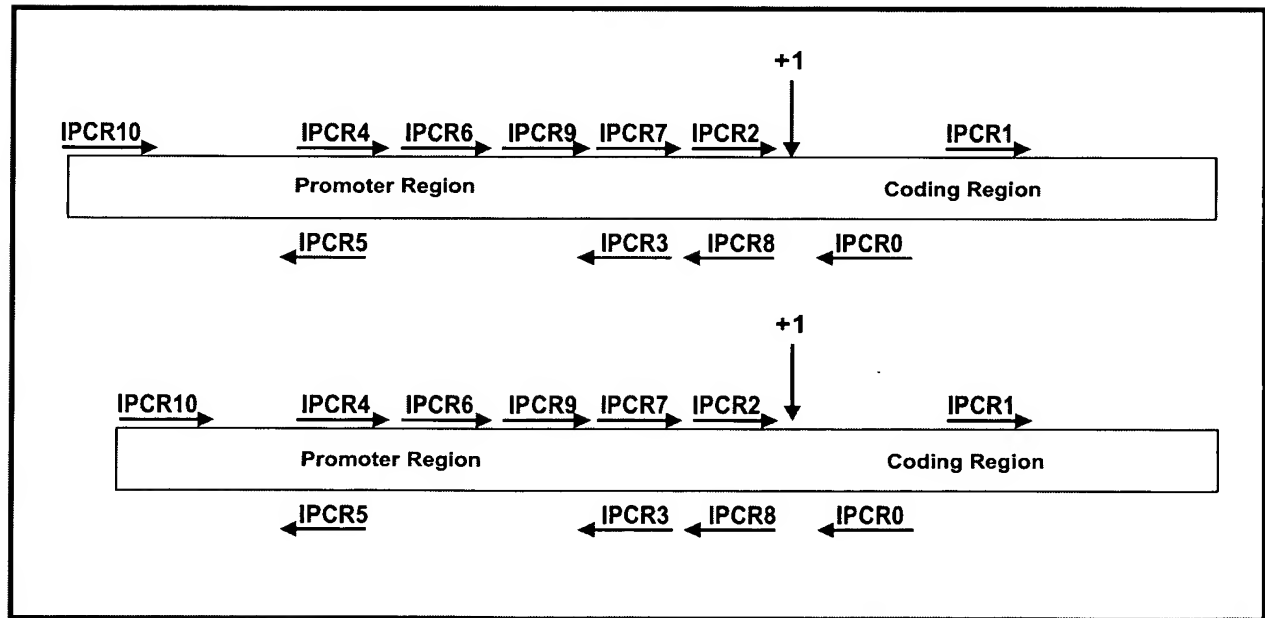


FIG. 23

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